

## SEQUENCE LISTING

- <110> ISHIWATA, TETSUYOSHI
- <110> SAKURADA, MIKIKO
- <110> KAWABATA, AYAKO
- <110> NAKAGAWA, SATOSHI
- <110> NISHI, TATSUNARI
- <110> KUGA, TETSURO
- <110> SAWADA, SHIGEMASA
- <110> TAKEI, MASAMI
- <110> SHIBATA, KENJI
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846

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Thr Lys Tyr Asp Val Leu Pro Tyr Ser Ile Arg Val Leu Leu Glu Ala
35 40 45

gct gta cga aat tgt gat ggc ttt tta atg aag aag gaa gat gtt atg 192 Ala Val Arg Asn Cys Asp Gly Phe Leu Met Lys Lys Glu Asp Val Met 50 55 60 aac att tta gac tgg aaa acc aaa caa agc aat gtt gaa gtg ccc ttt Asn Ile Leu Asp Trp Lys Thr Lys Gln Ser Asn Val Glu Val Pro Phe ttc cct gcc cgt gtt ctt ctt caa gat ttt act gga ata cca gca atg Phe Pro Ala Arg Val Leu Leu Gln Asp Phe Thr Gly Ile Pro Ala Met gtg gat ttt gct gct atg agg gag gca gtg aaa act ctt gga ggt gat Val Asp Phe Ala Ala Met Arg Glu Ala Val Lys Thr Leu Gly Gly Asp cct gag aaa gtc cat cct gct tgt ccg aca gat ctt aca gtt gac cat Pro Glu Lys Val His Pro Ala Cys Pro Thr Asp Leu Thr Val Asp His tct tta caa att gac ttc agt aaa tgt gca ata cag aat gca cca aat Ser Leu Gln Ile Asp Phe Ser Lys Cys Ala Ile Gln Asn Ala Pro Asn cct gga ggt ggt gac ctg cag aaa gca gga aag ctc tct cca ctt aaa Pro Gly Gly Asp Leu Gln Lys Ala Gly Lys Leu Ser Pro Leu Lys gtg cag cct aag aag ctt ccc tgc aga ggc cag act acc tgc cga gga Val Gln Pro Lys Lys Leu Pro Cys Arg Gly Gln Thr Thr Cys Arg Gly tct tgt gat tct gga gaa cta ggc cga aac tca gga aca ttt tct tcg Ser Cys Asp Ser Gly Glu Leu Gly Arg Asn Ser Gly Thr Phe Ser Ser cag att gag aat aca ccc atc ctg tgt cct ttt cat ttg caa cca gtg Gln Ile Glu Asn Thr Pro Ile Leu Cys Pro Phe His Leu Gln Pro Val cct gaa cct gaa aca gtg tta aaa aat caa gaa gta gaa ttc ggc aga Pro Glu Pro Glu Thr Val Leu Lys Asn Gln Glu Val Glu Phe Gly Arg aat cga gag agg ctt cag ttt ttt aag tgg agt tca aga gtt tta aag Asn Arg Glu Arg Leu Gln Phe Phe Lys Trp Ser Ser Arg Val Leu Lys aat gtg gca gtg atc cct cct gga act gga atg gct cat caa ata aac Asn Val Ala Val Ile Pro Pro Gly Thr Gly Met Ala His Gln Ile Asn

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tcctagtgac	aaaagttgag	aaactacctt	taaataagac	agtgaggtaa	caaatgt	357
<210> 24 <211> 219 <212> DNA <213> Homo	sapiens					
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aatagttaat	agctgtatta	gccagaaaat	ggtgtaagga	caacaggcta	actaaccctg	180
tcacttgtta	tgctaaaatt	aagtctagat	agagtcctc			219
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tccttacact	ggtgttcccc	aaagtgaggt	gaattgccag	ccactgggag	tcagggccag	120
ttacataaga	cattctcggt	aagccccctt	tgggtatccc	aaataaggac	tggggtgggt	180
ttatgtgtag	tccattatta	acaactaaac	gaacaaacct	agtgaattgc	aataaattca	240
caccaacaga	a					251
<210> 26 <211> 233 <212> DNA <213> Homo	sapiens					
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acaggatttc	taagaagtgg	aacagtctcc	aggggtgtgg	arctcatcgc	tcaaggcagg	120
ttatcttatc	tgaataattt	tgtctgttga	ctattgggat	agttctcctt	cagatgagct	180
gaaattttct	ccatagette	ctctattaaa	cccaattcca	cttctcaggg	tca	233

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<211> 176
<212> DNA
<213> Homo sapiens
<400> 27
caaaagcgct gaagttaagc attaatacgc cagattcatg atttatgatc agtatccaaa
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actccaacta caaacaatgc aaagtagtgc tcctcagtat tatttttgca attgttagta
                                                                      120
                                                                      176
atgttaagca tcaaggaaaa taaaacacat cattgcacat tacagccgca aaaaac
<210> 28
<211> 241
<212> DNA
<213> Homo sapiens
<400> 28
agagagtaaa gcaagctatt ttgacagcaa cctaataaca gctgtcttct tccacttctt
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ggctaactca tcccccagat agccttcttt tctcttatca attccctgtt gcaacaataa
                                                                       120
taaatgccac acctgatgga gtcattaggc actttcctag tgacaagtgc ctaggacaga
                                                                       180
ggagaaaaca aagaaacact gacaaccact gaaaactgac atatcaggcc aggcatgtca
                                                                       240
                                                                       241
C
<210> 29
<211> 217
<212> DNA
<213> Homo sapiens
<400> 29
gctggagagg tggtgatgtt gctgaataat tgctttttaa agctggaggg gacttccaag
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aggcaacact	ttgctcacaa	tcctacagat	ctacttcacc	tgtaactaca	attttcctga	180
agacatagaa	gaaaaatcaa	ttgttctaat	ccatatg			217
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cacacacgaa	gttcctagtt	cactgggact	tcctgatttg	ttcttttagc	ttgctccttc	120
tcacctagaa	gctctgttta	tttctgagca	accctggggc	ttgtctcata	ggacaggatt	180
tatttatctc	atcaaggctg	agtgtgcctt	aggaagtcat	aaacataaaa	aga	233
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actacctctc	: aggtcatgat	aaagaccggc	c cgggcagaaa	a cactgtaato	c ccagctactc	120
gggagcctga	a ggcatgagaa	ı tcacttgaad	c ctgggaggto	g gaggttgcca	a tgagccgaga	180
tcacgccatt	gcactacago	c cttggcgaca	a agagtgaaad	c tocatotg		228

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<210> 32

<211> 298

<212> DNA

<213> Homo sapiens

<400> 32

ategetttt etgaaatagg tateeettga tgtegactat ttgattteag eeagtegttt 120
etetetggea gtgeteeetg eaaatgtgte ettteaagaa aacaaaacet geaagtgget 180
tgtaatgtae eatgacetta teatgtgaag gacaaatgge tettgtgett attagatage 240
agatgaactg atgaactgaa ttettggtet gaagetttga taaggteaga tgtetttg 298

<210> 33

<211> 291

<212> DNA

<213> Homo sapiens

<400> 33

acttcgaagg gaaaaagagg aaggaaaagg actgttaata aaataacaaa ggcagcaatc 60
agaatgaacc agagccagga cagcgtaaag gctaggttca cagtgagatg aaagaacctg 120
aaaacaagtt taaaactcaa aagaggatta ttctcaagtt atactacagt gaaaaaacat 180
ggaaaaacac aaaaaggaca ggcaataagg cacaggcata catacaaggc aaattgtaac 240
acaatattta cttgcaaaag agcccacaga gacatgtcaa tgaagtcata g 291

<210> 34

<211> 230

<212> PRT

<213> Homo sapiens

<400> 34

Met Glu Asp Gly Phe Leu Asp Asp Gly Arg Gly Asp Gln Pro Leu His 

Ser Gly Leu Gly Ser Pro His Cys Phe Ser His Gln Asn Gly Glu Arg 

Val Glu Arg Tyr Ser Arg Lys Val Phe Val Gly Gly Leu Pro Pro Asp 

Ile Asp Glu Asp Glu Ile Thr Ala Ser Phe Arg Arg Phe Gly Pro Leu 

Ile Val Asp Trp Pro His Lys Ala Glu Ser Lys Ser Tyr Phe Pro Pro 

Lys Gly Tyr Ala Phe Leu Leu Phe Gln Asp Glu Ser Ser Val Gln Ala 

Leu Ile Asp Ala Cys Ile Glu Glu Asp Gly Lys Leu Tyr Leu Cys Val 

Ser Ser Pro Thr Ile Lys Asp Lys Pro Val Gln Ile Arg Pro Trp Asn 

Leu Ser Asp Ser Asp Phe Val Met Asp Gly Ser Gln Pro Leu Asp Pro 

Arg Lys Thr Ile Phe Val Gly Gly Val Pro Arg Pro Leu Arg Ala Val 

Glu Leu Ala Met Val Met Asp Arg Leu Tyr Gly Gly Val Cys Tyr Ala 

Gly Ile Asp Thr Asp Pro Glu Leu Lys Tyr Pro Lys Gly Ala Gly Arg 

Val Ala Phe Ser Asn Gln Gln Ser Tyr Ile Ala Ala Ile Ser Ala Arg 

Phe Val Gln Leu Gln His Gly Glu Ile Asp Lys Arg Val Ser Leu Ile 

Leu His Phe Gly Lys Phe 230 225

<210> 35

<211> 143

<212> PRT

<213> Homo sapiens

<400> 35

Met Gly Ser Asp Lys Arg Val Ser Arg Thr Glu Arg Ser Gly Arg Tyr 10 5 1

Gly Ser Ile Ile Asp Arg Asp Asp Arg Asp Glu Arg Glu Ser Arg Ser 25 20

Arg Arg Arg Asp Ser Asp Tyr Lys Arg Ser Ser Asp Asp Arg Gly 40 35

Asp Arg Tyr Asp Asp Tyr Arg Asp Tyr Asp Ser Pro Glu Arg Glu Arg 60 55 50

Glu Arg Arg Asn Ser Asp Arg Ser Glu Asp Gly Tyr His Ser Asp Gly 75 70 65

Asp Tyr Gly Glu His Asp Tyr Arg His Asp Ile Ser Asp Glu Arg Glu 90 85

Ser Lys Thr Ile Met Leu Arg Gly Leu Pro Ile Thr Ile Thr Glu Ser 110 105 100

Asp Ile Arg Glu Met Met Glu Ser Phe Glu Gly Pro Gln Pro Ala Asp 120 115

Val Arg Leu Met Lys Arg Lys Thr Gly Glu Ser Leu Leu Ser Ser 140 135 130

<210> 36

<211> 104

<212> PRT

<213> Homo sapiens

<400> 36

Met Pro His Met Leu Ser Gln Leu Ile Ala Gly Gly Val Ser Thr Ser 15 10 1

Cys Val Thr Ala Leu Gly Glu Glu Thr Gly Ala Trp Phe Pro Val Tyr 20 25 30

Leu Ser His Ala Ser Ser Pro Phe Ala Asp Leu Val Phe Cys Pro Phe 35 40 45

Ala Glu Ile Asn His Ser Gln Glu Tyr Asp Asn Met Arg Gly Pro Val 50 55 60

Ser Pro Pro Asn Lys Gln Phe Asn Leu Gly Val Ile Phe Gly Ile Pro 65 70 75 80

Asn Asn Cys Arg Phe Pro Thr Asp Asn Lys Ile Thr Glu Lys Gln Leu 85 90 95

Leu Gly Asn Val Leu Asn Tyr Pro 100

<210> 37

<211> 133

<212> PRT

<213> Homo sapiens

<400> 37

Met Asn His Pro Trp His Val Cys Phe Leu Phe Lys Val Leu Arg Tyr
1 5 10 15

Tyr Pro Thr Ala Pro Ile Leu Lys Trp Thr His Thr Val Ser Cys Ser 20 25 30

Trp Cys Arg Ser Val Leu Arg Glu Val Val Gly Asn Val Ser Leu Ser 35 40 45

Glu Asn Phe Thr Ile Ser Ala Phe Cys Pro Glu Leu Thr Pro Phe Pro 50 55 60

Asp Gln Gly Thr Ser Thr Met Ile Ser Phe Leu Glu Lys Phe Asn Lys
65 70 75 80

Ser Lys Arg Glu Arg Leu Glu Leu Met Leu His Phe Tyr Ser Val Leu 85 90 95

Ser Leu Glu Pro Ala Val Ala Glu His Trp Ser Gly Glu Phe Glu Lys 100 105 110 Trp Lys Val Gly Phe Phe His Pro Leu Lys Arg Glu Asp Gly Phe Phe 120 115

Thr Arg Thr Asp Ile 130

<210> 38

<211> 133

<212> PRT

<213> Homo sapiens

<400> 38

Met Asn His Pro Trp His Val Cys Phe Leu Phe Lys Val Leu Arg Tyr 10 5 1

Tyr Pro Thr Ala Pro Ile Leu Lys Trp Thr His Thr Val Ser Cys Ser 25 20

Trp Cys Arg Ser Val Leu Arg Glu Val Val Gly Asn Val Ser Leu Ser 40 35

Glu Asn Phe Thr Ile Ser Ala Phe Cys Pro Glu Leu Thr Pro Phe Pro 60 55 50

Asp Gln Gly Thr Ser Thr Met Ile Ser Phe Leu Glu Lys Phe Asn Lys 75 70 65

Ser Lys Arg Glu Arg Leu Glu Leu Met Leu His Phe Tyr Ser Val Leu 90 85

Ser Leu Glu Pro Ala Phe Ala Glu His Trp Ser Gly Glu Phe Glu Lys 110 105 100

Trp Lys Val Gly Phe Phe His Pro Leu Lys Arg Glu Asp Gly Phe Phe 125 120 115

Thr Arg Thr Asp Ile 130

<210> 39

<211> 128

<212> PRT

<213> Homo sapiens

<400> 39

Met Asp Ala Val Ala Val Tyr His Gly Lys Ile Ser Arg Glu Thr Gly 

Glu Lys Leu Leu Ala Thr Gly Leu Asp Gly Ser Tyr Leu Leu Arg 

Asp Ser Glu Ser Val Pro Gly Val Tyr Cys Leu Cys Val Leu Tyr His 

Gly Tyr Ile Tyr Thr Tyr Arg Val Ser Gln Thr Glu Thr Gly Ser Trp 

Ser Ala Glu Thr Ala Pro Gly Val His Lys Arg Tyr Phe Arg Lys Ile 

Lys Asn Leu Ile Ser Ala Phe Gln Lys Pro Asp Gln Gly Ile Val Ile 

Pro Leu Gln Tyr Pro Val Glu Lys Lys Ser Ser Ala Arg Ser Thr Gln 

Gly Thr Thr Gly Ile Arg Glu Asp Pro Asp Val Cys Leu Lys Ala Pro 

<210> 40

<211> 343

<212> PRT

<213> Homo sapiens

<400> 40

Met Asp Ala Pro Lys Ala Gly Tyr Ala Phe Glu Tyr Leu Ile Glu Thr 

Leu Asn Asp Ser Ser His Lys Lys Phe Phe Asp Val Ser Lys Leu Gly 

Thr Lys Tyr Asp Val Leu Pro Tyr Ser Ile Arg Val Leu Leu Glu Ala 

Ala Val Arg Asn Cys Asp Gly Phe Leu Met Lys Lys Glu Asp Val Met 

Asn Ile Leu Asp Trp Lys Thr Lys Gln Ser Asn Val Glu Val Pro Phe 

Phe Pro Ala Arg Val Leu Leu Gln Asp Phe Thr Gly Ile Pro Ala Met 85 90 95

Val Asp Phe Ala Ala Met Arg Glu Ala Val Lys Thr Leu Gly Gly Asp

100 105 110

Pro Glu Lys Val His Pro Ala Cys Pro Thr Asp Leu Thr Val Asp His

Ser Leu Gln Ile Asp Phe Ser Lys Cys Ala Ile Gln Asn Ala Pro Asn 130 135 140

Pro Gly Gly Gly Asp Leu Gln Lys Ala Gly Lys Leu Ser Pro Leu Lys 145 150 150

Val Gln Pro Lys Lys Leu Pro Cys Arg Gly Gln Thr Thr Cys Arg Gly 165 170 175

Ser Cys Asp Ser Gly Glu Leu Gly Arg Asn Ser Gly Thr Phe Ser Ser 180 185 190

Gln Ile Glu Asn Thr Pro Ile Leu Cys Pro Phe His Leu Gln Pro Val 195 200 205

Pro Glu Pro Glu Thr Val Leu Lys Asn Gln Glu Val Glu Phe Gly Arg 210 215 220

Asn Arg Glu Arg Leu Gln Phe Phe Lys Trp Ser Ser Arg Val Leu Lys 225 230 235 240

Asn Val Ala Val Ile Pro Pro Gly Thr Gly Met Ala His Gln Ile Asn 245 250 255

Leu Glu Tyr Leu Ser Arg Val Val Phe Glu Glu Lys Asp Leu Leu Phe 260 265 270

Pro Asp Ser Val Val Gly Thr Asp Ser His Ile Thr Met Val Asn Gly 275 280 285

Leu Gly Ile Leu Gly Trp Gly Val Gly Gly Ile Glu Thr Glu Ala Val 290 295 300

Met Leu Gly Leu Pro Val Ser Leu Thr Leu Pro Glu Val Val Gly Cys 305 310 315 320

Glu Leu Thr Gly Ser Ser Asn Pro Phe Val Thr Ser Ile Asp Val Val

325 330 335

Leu Gly Ile Thr Lys Val Ser 340

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<211> 305

<212> DNA

<213> Homo sapiens

<400> 41

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attcatagac ataaaccctc attttaatta gtggatctgg atttttgtca tatgtggaat 120
cataatttaa acaaaatcaa ctaagatgat ccaagttcca cacaactgca cttcaatatt 180
caagtcggtg tgaagatgcc tgactactgc gtcacaagat tctgagctgt cgtaaaaagc 240
ctggctcgtg gtttctattt atagtgtaca catgttgggt tataatcaca aacctggaac 300
tctgt

<210> 42

<211> 256

<212> DNA

<213> Homo sapiens

<400> 42

gaaaccacgg cttacaccta gagacagcat tcagatatag acgggatact tgtgttagtc 60
agttccttta taacaggtga atctctctcc cactgcttca acactgcgtg acaaagccaa 120
ttgggaagca gctttacaaa tgtgacttga cttggggatc ttcttgatac tttgccatgg 180

caaggaacaa gccgcctgaa	ctaaatgcca	ctccatttga	ttccacgctt	aaagtaacca	240
tgcaaccgac tatagt					256
<210> 43 <211> 244 <212> DNA <213> Homo sapiens					
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aattccttcc ccactggggg	aaagcaaatc	atcaggccca	ttgcaaaaac	tgctcttggt	120
tgagetteet geettaaate	atacccacag	tgaatggcgt	ccctttatca	ccgctaatga	180
ctctgacatc tctctccact	cacatgtgag	cctcctcagc	tctcganaaa	caagtengte	240
tcgg					244
<210> 44 <211> 258 <212> DNA <213> Homo sapiens					
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tactctccaa tggtgatgaa	gggagatgtc	tgggggaaat	ccagcaggat	gttgatttag	120
tatgtacaca gtgagaggat	acttgtagag	aacctagaat	cttctctgaa	tgtgacgggc	180
cctcagagat aattgttaac	agataagtgg	atgattaaat	acacttcctc	cagtaggcta	240

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258
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<211> 26
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<213> Artificial Sequence
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<400> 45
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gggcttaata ttattcatag atcgag
<210> 46
<211> 26
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 46
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gttattatac tatcaagtaa cccaac
<210> 47
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 47
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gtggatctgg atttttgtca tatgt
<210> 48
<211> 25
<212> DNA
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<213> Artificial Sequence

<220> <223> Description of Artificial Sequence: Synthetic DNA	
<400> 48 gtttgtgatt ataacccaac atgtg	25
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<220> <223> Description of Artificial Sequence:Synthetic DNA	
<400> 49 gaaggggaag agacattaaa ttatc	25
<210> 50 <211> 24 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Synthetic DNA	
<400> 50 gcttctaaat ctcctgagtc actt	24
<210> 51 <211> 24 <212> DNA <213> Artificial Sequence	
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<400> 51 gacaatgagt aagaagaag aggg	24

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<210> 52
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 52
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gtccagtccc ttggtttatt tgtc
<210> 53
<211> 25
<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 53
ggtacccagt ttcaaattaa catgg
                                                                        25
<210> 54
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 54
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gattetteaa etgeeaaact tgtte
<210> 55
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
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<400> 55
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gctgatgctt ttctatctga cttc
<210> 56
<211> 22
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<220>
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<400> 56
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gaccaggact gaacagaggt ga
<210> 57
<211> 25
<212> DNA
<213> Artificial Sequence
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<400> 57
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gcttatagac catgtttgta gtagg
<210> 58
<211> 25
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 58
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qtqaacaaat qctaaatcag acatg
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<210> 59 <211> 22

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<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 59
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gccacgggtt tcccatatcg aa
<210> 60
<211> 24
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 60
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gactatactt aggaacctct gcaa
<210> 61
<211> 24
<212> DNA
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<400> 61
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gttctgctct cagcagattg gtta
<210> 62
<211> 24
<212> DNA
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 62
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gccaacatct gaactaaata ctgc	24
<210> 63 <211> 25 <212> DNA <213> Artificial Sequence	
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<400> 63 gttcagtgaa tgttacctag aaaca	25
<210> 64 <211> 24 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence:Synthetic DNA	
<400> 64 ggagtgaaaa ctgtcttgtt catc	24
<210> 65 <211> 25 <212> DNA <213> Artificial Sequence	
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<400> 65 gtatgacaaa tagtttctgc ctgat	25
<210> 66 <211> 25	

<212> DNA

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<400> 66
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gattaacaaa gatgtacaga ctgag
<210> 67
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 67
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gagacagcat tcagatatag acgg
<210> 68
<211> 22
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 68
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gcgtggaatc aaatggagtg gc
<210> 69
<211> 24
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic DNA
<400> 69
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gatggcctgt gtgaacagat taat
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<210> 70
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 70
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gagagatg tcagagtcat tagc
<210> 71
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 71
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gatccccaca atttcttgtg attg
<210> 72
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 72
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gttcccctaa aataatgtgg taatg
<210> 73
<211> 23
<212> DNA
<213> Artificial Sequence
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<220> <223> Description of Artificial Sequence: Synthetic DNA	
<400> 73 gaggatactc tccaatggtg atg	23
<210> 74 <211> 24 <212> DNA <213> Artificial Sequence	
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<400> 74 gtcttaacat ctagcctact ggag	24
<210> 75 <211> 24 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Synthetic DNA	
<400> 75 gagaggagcc atgtatacaa acca	24
<210> 76 <211> 26 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Synthetic DNA	
<400> 76 gcacgcagga tcagatatag taattc	26

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<210> 77
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 77
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gctgaaacct aagctgaagg aagg
<210> 78
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 78
                                                                         22
gtccctcacc tcagatcaca cc
<210> 79
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 79
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gctatctacc tggcaggaaa agag
<210> 80
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
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qaqtttctta ctatgatctg gattc
<210> 81
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 81
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gcaaaatgta ctcagcttca atcac
<210> 82
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 82
gtaaatgcag tactgttctg atcc
                                                                         24
<210> 83
<211> 26
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 83
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gaatgcttca ttctcattgt ttaagg
<210> 84
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<211> 24

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<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 84
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gtcactagga ttccacagaa cttc
<210> 85
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 85
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gaggtagggc ttcccttcgc ta
<210> 86
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 86
                                                                         25
gcataacaag tgacagggtt agtta
<210> 87
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 87
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22
ggtgctcctt ccttacactg gt
<210> 88
<211> 23
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic DNA
<400> 88
                                                                        23
qactacacat aaacccaccc cag
<210> 89
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 89
                                                                         24
gggtacagga tttctaagaa gtgg
<210> 90
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic DNA
<400> 90
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ggagaaaatt tcagctcatc tgaag
<210> 91
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<211> 24 <212> DNA

<213> Artificial Sequence	
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<400> 91 gctgaagtta agcattaata cgcc	24
<210> 92 <211> 23 <212> DNA <213> Artificial Sequence	
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<400> 92 gcggctgtaa tgtgcaatga tgt	23
<210> 93	
<211> 24 <212> DNA	
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gaggggactt ccaagagtct ct
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gtcttcagga aaattgtagt tacag
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gtttaactac ctctcaggtc atga
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gtcgccaagg ctgtagtgca at
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<400> 101
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gaaataggta tcccttgatg tcga
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gaccaagaat tcagttcatc agtt
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gaatgaacca gagccaggac ag
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gccttgtatg tatgcctgtg cc
<210> 105
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gtttttttt ttttta
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<210> 109 <211> 17

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cagagtgatg gatatcaa
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atgaaagtgc cagtgtgcca tg
<210> 112
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cccatcacca tcttccagga gc

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<210> 113
<211> 26
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ttcaccacct tcttgatgtc atcata
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<400> 114
Cys Pro Leu Lys Arg Glu Asp Gly Phe Phe Thr Arg Thr Asp Ile
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                                                           15
  1
<210> 115
<211> 16
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<221> MOD RES
<222> (16)
<223> AMIDATION, GluAmide
<400> 115
Cys Ser Phe Leu Glu Lys Phe Asn Lys Ser Lys Arg Glu Arg Leu Xaa
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  1
                  5
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<210> 116

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<211> 15
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<222> (15)
<223> AMIDATION, GlyAmide
<400> 116
Cys Ala Glu His Trp Ser Gly Glu Phe Glu Lys Trp Lys Val Xaa
                                                           15
                                      10
<210> 117
<211> 16
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<400> 117
Cys Glu Ile Asp Lys Arg Val Ser Leu Ile Leu His Phe Gly Lys Phe
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<210> 118
<211> 15
<212> PRT
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Cys Arg Leu Met Lys Arg Lys Thr Gly Glu Ser Leu Leu Ser Ser
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  1
                  5
                                      10
<210> 119
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<223> Description of Artificial Sequence: Synthetic Peptide
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Cys Thr Ser Ile Asp Val Val Leu Gly Ile Thr Lys Val Ser
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                  5
<210> 120
<211> 16
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<223> AMIDATION, LysAmide
<400> 120
Cys Ser Ala Glu Thr Ala Pro Gly Val His Lys Arg Tyr Phe Arg Xaa
                                                           15
                                      10
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<210> 121
<211> 16
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic Peptide
<400> 121
Cys Lys Ile Thr Glu Lys Gln Leu Leu Gly Asn Val Leu Asn Tyr Pro
                                                           15
                                      10
```

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